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Steven PELECH

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The benefit of the filing date of the following priority foreign application(s) in the following foreign country is hereby requested, and the right of priority provided in 35 U.S.C. § 119 is hereby claimed.

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Patent Application No(s).: 2,290,204 and 2,290,335

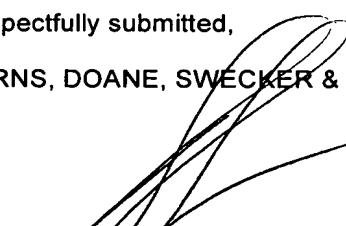
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In support of this claim, enclosed is a certified copy(ies) of said foreign application(s). Said prior foreign application(s) is referred to in the oath or declaration. Acknowledgment of receipt of the certified copy(ies) is requested.

Respectfully submitted,

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Specification and Drawings, as originally filed, with Application for Patent Serial No:
2,290,335, on November 19, 1999, by THE UNIVERSITY OF BRITISH COLUMBIA,
assignee of Steven Pelech, for "Multiblot Kinase Analysis".

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Agent certificateur/Certifying Officer

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Canada

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MULTIBLOT KINASE ANALYSIS

Background of the Invention

5 The human genome is believed to contain about 120,000 genes, which are present in each of the 50 trillion nucleated cells of the body. At any given moment in each cell, however, only about 20% of these genes are actively transcribed for the production of the proteins that they encode. The unique complement of proteins that each cell expresses is referred to as its "proteome". It is critical that the levels of expression and activity of the 10 proteins in a cell are tightly regulated. This is achieved through a subset of about 10% of these proteins, which are dedicated to cell communications and control. One of the largest classes of proteins involved in cell signaling are enzymes called protein kinases. Protein kinases control other proteins by catalyzing their phosphorylation, which is a process that can be reversed by protein phosphatases. Often protein kinases operate within signalling 15 pathways that are further integrated into networks. The unique complement of protein kinases expressed in a cell is referred to herein as its "kineome".

There are different amino acids that can be phosphorylated by protein kinases. Most commonly, this occurs on serine and threonine, and to a much lesser extent on tyrosine. For example, in skin cells called fibroblasts, 90% of measurable protein kinases catalyze the 20 serine and threonine phosphorylation of proteins, whereas a different class of related enzymes generally carry out tyrosine phosphorylation. Therefore, protein-serine/threonine kinases are responsible for most protein phosphorylation events in cells. For convenience, these kinases are referred to herein as protein-serine kinases, although they also phosphorylate proteins on threonine.

25 Approximately fifty of the hundred or so known genes that have been directly linked to induction of cancer (i.e. oncogenes) encode protein kinases. The remainder of the oncogenes specify proteins that either activate kinases or are phosphorylated by kinases. Most of the oncogene-encoded protein kinases are tyrosine-specific, but several are protein-serine kinases such as protein kinase C, Raf1, Akt, ILK-1, Tpl2, and Mos. Although 30 the findings are less direct, aberrant cell signalling through protein kinases has also been linked to cardiovascular disease, diabetes, inflammation, arthritis and other immune disorders, and neurological disorders such as Alzheimer's disease. Over 400 human diseases have been linked to defective signalling through protein kinases.

Essentially all signalling proteins, if they are not already protein kinases, appear to be regulators of protein kinases or their substrates. As signal transduction networks govern and co-ordinate all cellular functions, including cell structure, metabolism, reproduction, adaptation, differentiation and death, knowledge of the structure of signalling networks will 5 permit a complete understanding of how the cell operates under a diversity of conditions. Kineome analysis represents a key step in this process.

Kineome analysis will yield many practical benefits. The presence and state of activity of diverse protein kinases and their pathways are indicators of how a cell perceives its internal and external environments and how it is responding. Therefore, by monitoring the 10 kineome, it will be feasible to obtain a molecular diagnosis of a disease condition. Moreover, by inhibiting or activating the appropriate protein kinases by pharmacological intervention, antisense or gene therapy, it would be possible to "reprogram" the kineome to better treat the disease condition. For example, in cancer, the gain of function of one of over fifty different oncogene-encoded protein kinases may be pivotal for neoplastic transformation of cells. 15 Inhibition of the appropriate kinase or its downstream effectors could block the improper proliferative signalling and initiate apoptotic processes leading to programmed death of the tumor cells.

The genome sequencing projects for man, mouse and other organisms will permit the rapid identification of all of the protein kinases within the next few years. Elucidation of the 20 connections between these protein kinases in different cells will allow kineome analysis to then reach its full potential. Over two thousand different protein kinases are thought to be encoded by the human genome and several hundred are likely to be expressed within any given cell. All of these protein kinases will have to be tracked for a complete elucidation of the architecture of kinase networks. This is feasible through the employment of kinase 25 specific probes.

Most eukaryotic protein kinases are evolutionarily related, i.e. the genes of almost all protein-serine/threonine and protein-tyrosine kinases display sequence identity. In particular, there are 16 amino acid residues located in 10 subdomains in the catalytic region of protein kinases that are highly conserved. These amino acids allow unambiguous identification of 30 novel protein kinases following analysis of the primary structures of proteins as revealed by the nucleotide sequences of genes. With the complete sequencing of the genomes of humans

and other species, it will be possible to readily identify most if not all of the protein kinases.

From knowledge of the primary structure of a protein, it is feasible to produce nucleotide or antibody probes that are specific for that protein. Antibodies can be generated against the full length-expressed protein or portions. An effective strategy is to identify a 5 region of about 10 to 20 amino acids that are extremely well conserved in that protein in diverse species, but which does not appear in other proteins. Antibodies generated, for example in a rabbit or mouse, against a synthetic peptide based on this amino acid sequence will cross-react with the full-length protein that contains this sequence, with little or no cross-reactivity with other proteins. With the knowledge of the primary structures of all the protein 10 kinases and other proteins encoded by mammalian genomes, specific antigen peptides can be designed to elicit the production of antibodies against all of the protein kinases.

Most protein kinases appear to be activated as a consequence of their own phosphorylation by upstream kinases or by self-phosphorylation (termed autophosphorylation). Phosphorylation cannot be monitored by nucleic acid-based 15 approaches. However, phosphorylation of a protein can produce marked changes in its mobility on electrophoresis gels which are designed to act as a molecular sieve. Of such techniques, sodium dodecyl sulphate (SDS) - polyacrylamide gel electrophoresis (SDS-PAGE) has become the standard method for separation of proteins on the basis of their size for analytical and preparative purposes. This technique relies of the sieving effect of the 20 gel when proteins coated with negatively-charged detergent (eg. SDS) are drawn through the gel in an electric field. Smaller sized proteins are able to migrate through the gel faster than larger sized proteins. Proteins that differ by as little as a few hundred Daltons can be resolved by this method. Protein staining methods permit the visualization of discreet 25 proteins in the gel as individual bands in a bar code like pattern. When these proteins are transferred from the gel onto a nitrocellulose membrane, the locations of specific proteins can be identified with antibodies in an immunoblotting procedure referred to as Western blotting (see: Towbin; U.S. Patent No. 4,452,901).

Most proteomic analytical methods are based on two dimensional (2D) gel 30 electrophoresis by the standard method of Dr. Patrick O'Farrell described nearly 20 years ago. The 2D gel technique initially involves the separation of proteins in a first dimension based on their intrinsic charge in a pH gradient within a isoelectric focusing gel (typically a

tube gel). Proteins migrate through the isoelectric focusing gel in the presence of an electric field until they encounter a pH at which the protein no longer possesses an electric charge. This pH is known as the isoelectric point of a protein, and it is a distinguishing characteristic. Following electrophoresis in the first dimension, the isoelectric focusing gel is applied 5 length-wise to the top of a molecular sieve gel such as a SDS-PAGE gel, and electrophoresis is continued into the second dimension. When the 2D gel is stained with sensitive-dyes (eg. based on silver reagent), the various proteins inside a cell can be visualized as resolved spots. The greater amount of a given protein within a cell sample, the larger and darker its specific spot appears. Several thousand proteins can be distinguished from one and another 10 by this technique. If the protein samples have been obtained from cells that have been incubated with radioactive ^{32}P -phosphate, then the 2D gel can be exposed to x-ray film, and the phosphoproteins can be specifically detected. The more that a protein is phosphorylated or prevalent, the larger and more intense the spot on the x-ray film. The silver-staining of a 15 2D gel can be used to track the expression of proteins and their covalent modification by phosphorylation.

Public databases have been created that allow the identification of over a thousand different proteins on 2D gel proteomic maps. However, the positions of scarcely more than a dozen protein kinases are available. This reflects the fact that protein kinases are present at very minute levels in cells, and are often undetectable by even such sensitive protein dyes as 20 silver-stain. Typically, transduction proteins are expressed at a hundred- to a thousand-fold lower levels than structural proteins and metabolic pathway enzymes. Therefore, it has often been necessary to incorporate selective enrichment techniques such as antibody-based purification as a preliminary step prior to 2D gel electrophoresis.

Proteomic analysis of kinases has evolved to date in much the same fashion as 25 techniques for resolution of other proteins. One dimensional immunoblot techniques have been employed in which proteins from a smooth muscle homogenate are separated on an SDS-PAGE gel and then subjected to Western blotting followed by detection using a single polyclonal or monoclonal (MAB) anti-kinase antibody per blot (see: H. Togashi *et al.* (1997) "Quantitative Immunoblot Analysis of PKC Isoforms Expressed in Airway Smooth Muscle" 30 Am. J. Physiol. 272 (Lung Cell. Mol. Physiol. 16): L603-L607). Conventional wisdom is that simultaneous use of multiple antibodies on a single immunoblot necessitates the use of

MAB's (see: Coates, S.R. *et al.*; EP 025384 published January 27, 1988). Furthermore, 2D gel electrophoresis is thought to be the preferable technique for resolving complex protein mixtures. For example, Sanchez, J.C. *et al.* report the use of a mixture of nine MAB's for detection of different proteins indicative of oncogene expression (including the kinase 5 MEK-1) on a single immunoblot produced from 2D electrophoresis ("Simultaneous Analysis of Cyclin and Oncogene Expression Using Multiple Monoclonal Antibody Immunoblots" (1997) Electrophoresis 18:638-641).

While it is possible to visualize some protein kinases on 2D gels by immunoblotting techniques, we have determined that in most cases, a maximum of only four or five protein 10 kinases can be detected at a time by Western blotting of 2D gels with mixtures of protein kinase-specific antibodies. Furthermore, we have found that recovery of most protein kinases from a first dimension pH gradient gel, is less than 10 %. That means that 90 % or more of the protein kinases do not enter the second dimension gel and are therefore unresolved.

15 **Summary of Invention**

This invention provides a method for detection of multiple kinases or multiple kinase substrates, whereby the presence and phosphorylation state of a large number kinases and/or kinase substrate proteins may be tracked in a single sample. This method comprises:

20 (a) obtaining a sample to be tested for kinase or kinase substrate content;

(b) optionally performing one or more of:

25 (i) addition of at least one protein phosphatase to dephosphorylate proteins in the sample;

(ii) inactivating protein phosphatase in the sample;

30 (iii) addition of at least one kinase and ATP to the sample; and

- (iv) inactivating protein kinase in the sample.
- (c) performing SDS-PAGE on the sample to produce a pattern of separated kinase or kinase substrate moieties from the sample;
- 5 (d) transferring the pattern to a membrane;
- (e) treating the membrane with a panel of anti-kinase or anti-kinase substrate antibodies; and
- 10 (f) detecting the presence of antibodies from the panel bound to kinases or kinase substrates on the membrane.

When the method of this invention is employed to detect kinases in a sample either to elucidate the kinase profile of a tissue or cell type or to identify novel kinases, it is preferable that the panel of anti-kinase antibodies comprise polyclonal antibodies rather than MAB's. This departure from conventional wisdom increases the likelihood that new kinase proteins will be detected in the sample since polyclonal anti-kinase antibodies generally exhibit greater cross-reactivity to kinases as compared to anti-kinase MAB's. Despite the use of polyclonal antibodies, the panel may comprise from 2 to about 100 antibodies.

While (c) and (d) above in combination, is similar to standard Western blotting procedure, it is preferable that the electrophoresis gel be constructed to increase the likelihood that proteins will exhibit "band shift" between phosphorylated and dephosphorylated states. Typically, a protein will display reduced migration during SDS-PAGE when the protein is in a phosphorylated state. The reduced mobility may be as much as 1-5 kDa and this separation is enhanced by using a gel with a higher than normal acrylamide content and a lower than normal bisacrylamide content. While content of these gel components is normally adjusted to suite electrophoretic conditions and the average size of proteins to be separated, a 12.5% acrylamide/0.4% bisacrylamide gel is often suitable to achieve separation of phosphorylated and dephosphorylated kinases.

Brief Description of the Drawings

Figure 1, shows gels presenting comparison of the multi-kinase immunoblotting patterns of different rat tissues. Electrophoresis of detergent solubilized lysates prepared from rat brain (A), heart (B) and skeletal (C) muscle was performed, and the positions of various protein kinases was visualized by ECL detection. Protein kinases of smaller size migrated correspondingly closer to the bottom of the SDS-PAGE gel. Each of the 18 strips derived from each SDS-PAGE gel were probed with different panels of protein kinase antibodies.

Figure 2, shows a gel demonstrating effects of anti-IgM treatment for 5 min on protein kinases in the human Ramos B cell line. Electrophoresis of detergent solubilized lysates prepared from Ramos cells untreated (-) or exposed (+) to anti-IgM polyclonal antibody for 5 min was performed in alternating lanes, and the positions of various protein kinases was visualized by ECL detection. Each of the 14 paired strips derived from two SDS-PAGE gels were probed with different panels of protein kinase antibodies.

Figure 3A. Fig. 3A are gels showing differential effects of kinase inhibitors on band shifting of selected protein kinases. Electrophoresis of detergent solubilized lysates prepared from human ovarian surface epithelial cells were untreated (Lane 1) or exposed to 20 ng/ml of human hepatocyte growth factor (HGF) in the absence (Lane 2) or presence of PD98059 (Lane 3), SB203580 (Lane 4), LY294002 (Lane 5) or rapamycin (Lane 6) was performed. The effects of these treatments on the positions of Erk1 and Erk2 (Panel A), p38 Hog MAP kinase (Panel B), PKB1 (Panel C), PKB2 (Panel D) and S6 kinase (Panel E) as visualized by ECL detection are shown. The phosphorylated and band shifted forms of these kinases are denoted with a "p" before their name. The partial structures of the protein kinases pathways in which these enzymes operate and the known sites of action of these drugs are shown in Fig. 3B.

Figure 4, are gels showing detection of known kinases and putative kinases in normal and tumor breast tumour biopsy samples of four human patients. Detergent solubilized lysates prepared from tumour (T) and adjacent control (C) breast tissue were subjected to multi-

kinase profiling. In the left (A) panels, the increased levels of p38 MAP kinase, protein kinase B- α (PKB α), casein kinase 2 (CK2), protein kinase G (PKG) and cyclin-dependent kinase 8 (Cdk8) in the tumour samples is evident. Five of 12 proteins that were demonstrated to be elevated in tumours and not yet known for their identity are shown on the right (B) in
5 Figure 4.

Figure 5 is a Western Blot showing separation of Erk1, Erk2 and protein kinase C- β by 2D gel electrophoresis. Detergent solubilized rat brain extract (1 mg protein) was subjected to isoelectric focusing and SDS-PAGE. In the left most lane, 200 μ g of the brain extract was
10 directly applied to the same SDS-PAGE gel. Following 2D gel electrophoresis, the proteins were transferred to a nitrocellulose membrane, which was probed with antibodies for Erk1, Erk2 and PKC- β .

Detailed Description

15 In this specification, the term kinase refers to those members of the class of enzymes that catalyze a chemical reaction in which a phosphate group is transferred from adenosine triphosphate (ATP) to a recipient protein. In the content of this specification, such a recipient protein is termed a "kinase substrate". This chemical reaction is called
20 protein phosphorylation and is a reversible process with dephosphorylation being catalyzed by enzymes known as protein phosphatases. Kinases may be found in all organisms. In this specification, testing for kinase or kinase substrate content means determining the presence of at least one kinase or kinase substrate in a sample. Preferably the phosphorylation state of the kinase or kinase substrate will also be determined. In this
25 specification, a kinase or kinase substrate moiety is a protein having the characteristics of a kinase or kinase substrate which occupies a single position after electrophoresis in a SDS-PAGE gel.

30 Samples used in the method of this invention may be any cell or tissue homogenate, extract or other such sample which has been processed to purify or partially purify kinases or kinase substrates in the sample. This invention is particularly suitable for testing patient biopsy samples. Such samples may be manipulated to increase prevalence of desired cell

types in the sample. The sample will be typically prepared for electrophoresis using standard techniques, employing appropriate buffers which may contain various inhibitors or enzymes. For example, protease inhibitors may be present to reduce protein degradation on the sample. Where the sample is to be tested for kinase substrate content, it may be 5 desirable to add one or more protein phosphatases to dephosphorylate substrates which may already exist in a phosphorylated state in the sample. The protein phosphatase will then be inactivated with an appropriate phosphatase inhibitor (eg. β -glycerophosphate, sodium fluoride or sodium orthovanadate) and a selected protein kinase or mixture of protein kinases is then added to the sample to phosphorylate those substrates present which are 10 specific to kinase added to the sample. Alternatively, endogenous kinases in the sample may be relied upon to phosphorylate dephosphorylated substrates in the sample.

SDS-PAGE employed in this invention is gel electrophoresis performed in a single dimension, typically using a slab shaped gel or a series of tube gels. The gel may be constructed and used employing standard methods, electrophoresis buffers and 15 electrophoresis equipment. The gel may comprise a stacking gel and a separation gel. Commercial kits and equipment are available for performing SDS-PAGE. Preferable contents of the separation gel range from 10% to 15% (acrylamide) and 0.2 to 2% (bisacrylamide). For separation of kinases, an electric current will typically be applied to the gel until proteins with a molecular mass of less than about 25-27 kDa are eluted from 20 the bottom of the gel as protein kinases do not have a molecular mass less than the latter amount. Once electrophoresis is complete resulting in a pattern of separated protein moieties in the gel, the pattern is transferred to any membrane (eg. nitrocellulose, PVDF, nylon, etc.) that is suitable for use in the Western Blotting technique. Transfer is typically done by standard electro-transfer techniques. Once the pattern is transferred to the 25 membrane, the membrane may be cut into strips each of which will typically contain a pattern separated from a single sample (eg. a test sample or a control sample).

Once the electrophoresis and Western blotting aspects of this invention are complete, the resulting membrane or membrane strips are probed with a panel of different antibodies that react with distinct categories, subsets, isoforms, etc. of protein kinases or 30 kinase substrates. The panel may be applied in one step as a mixture of antibodies or, the antibodies may be applied sequentially to the membrane. Binding of such antibodies to

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moieties present on the membrane is then detected using any suitable immunoassay procedure (eg. see: Stites and Terr (eds) "Basic and Clinical Immunology", (7 ed) 1991). A particularly suitable procedure is to treat the antibodies in the panel as primary antibodies in a "sandwich" type assay. Unbound primary antibodies are washed away or otherwise 5 removed. The membrane is then treated with secondary antibodies which are reactive with the primary antibodies. The secondary antibody may be bound to a detectable label or fused with an enzyme. Secondary antibody bound to primary antibody is detected by observing the label or the activity of the fused enzyme. Suitable labels and enzymes are known in the art and include magnetic or coloured beads, fluorescent dyes, radiolabels. 10 horseradish peroxidase, alkaline phosphatase, etc. The enzyme linked sandwich type assay (ELISA) is a particularly suitable methodology for use in this invention.

Antibodies for use in this invention may be obtained commercially or prepared using standard techniques. A variety of anti-kinase and anti-kinase substrate polyclonal antibodies are commercially available from various sources, including the following:

15

Biomol Research Laboratories, Inc. (Plymouth Meeting, Pennsylvania)

Biosource International, Inc. (Camarillo, California)

Promega Corporation (Madison, Wisconsin)

Santa Cruz Biotechnology (Santa Cruz, California)

20

Sigma (Saint Louis, Missouri)

StressGen Biotechnologies Corp. (Victoria, British Columbia)

Transduction Laboratories (Lexington, Kentucky)

Upstate Biotechnology Inc. (Lake Placid, New York)

Zymed Laboratories Inc. (South San Francisco, California)

25

Antibodies to new kinases may be prepared as described below. Typically, new kinases are partially purified by techniques such as column chromatography and SDS-PAGE. Microsequencing of partially purified kinases permits comparison to known kinases and possible development of immunological techniques for recovery of more of the 30 new kinase by making use of cross reactivity with known antibodies. Antibodies can be raised against protein kinases or substrates in various host animals, including but not

limited to cattle, horses, rabbits, goats, sheep and mice. Polyclonal antibodies can be obtained from immunized animals and tested for specificity using standard techniques. Alternatively, monoclonal antibodies may be prepared using any technique that provides for production of antibody molecules by continuous cell lines in culture, including the 5 hybridoma technique of Kohler and Millstein, the human B-cell hybridoma technique, and the EBV-hybridoma technique. Alternatively, techniques for the production of single chain antibodies and antibody fragments that contain specific binding sites for a protein kinase or substrate may be generated by known techniques and employed in this invention. Such fragments include $F(ab')_2$ fragments that may be generated by digestion of an intact 10 antibody molecule and Fab fragments that may be generated by severing disulfide bridges in $F(ab')_2$ fragments or through the use of Fab expression libraries.

Preferably, none of the antibodies in a given mixture to be used as a panel in this invention will cross-react with proteins that overlap in size. This may compromise interpretation of the experimental findings. Each antibody panel mixture should be blended 15 to avoid such overlaps. Furthermore, every mixture should be adjusted for the concentration of each antibody so that there is optimal detection of the individual target kinases in diverse cell and tissue samples.

Following incubation of the strips with different mixtures of primary antibodies, the strips are incubated with a secondary antibody (eg. a goat antibody that recognizes rabbit 20 antibody) that reacts with the primary antibody. The secondary antibody is fused with an enzyme (eg. alkaline phosphatase or horse radish peroxidase) to facilitate detection of the positions of the primary antibody, to which it binds by producing a light emission in an enzymatic reaction. The separate strips may be reassembled to appear in the order of the original membrane. The reassembled membrane may be subjected to enhanced 25 chemiluminescence (ECL) and exposure to x-ray film or detected by a phosphoimager (eg. Fluor-S Max Multi-imager from Bio-Rad Laboratories). In this indirect manner, the original positions of resolved protein kinases can be visualized as dark bands on a transparent background. The intensity of the bands can be quantitated by densitometric analysis. In many cases, quantitation of the amounts of a given protein kinase in the 30 upper, phosphorylated form and the lower dephosphorylated form can provide an accurate measurement of how much of the kinase is in the inactive and active states.

This invention offers advantages over standard 2D gel proteomic methods. This technique can be applied to any cell or tissue sample. No prelabelling with radioisotopes is necessary, because kinase detection is based on immunoreactivity. The technology could
5 be adapted for wide scale diagnostic applications because the patterns of protein kinase expression are stable for periods of up to six hours before an organ is subjected to fractionation and freezing, providing the organ is stored during this time over ice. This procedure can be carried out within two days from start to finish. By contrast, the 2D gel electrophoresis approach is extremely laborious, much more difficult to render and takes at
10 least twice the time. This invention provides the ability to compare multiple samples side by side. Whereas two or more samples can be analyzed on the same 1D gel, a 2D gel can only be used for a single sample. It is more difficult to compare two different samples by the 2D gel route, because of potential variations in the setting up, running and analysis of separate 2D gels.

15 One of the reasons why 2D gel electrophoresis has become the industry standard for proteomic analysis is the remarkable resolving power of the method and potentially thousands of spots can be distinguished on a 2D gel. Most of these spots, however, are "fuzzy" in appearance and may be overlapping. The method of this invention provides much tighter protein bands with a 2- to 4-fold better resolution in the SDS-PAGE
20 size-separation dimension. With detection based on immunoreactivity, the background of metabolic enzymes and structural proteins is essentially eliminated. This background is problematic even for 2D gel maps of phosphoproteins, since a third of all the proteins inside of cells appear to be phosphorylatable.

In one exemplary embodiment of this invention, about 50 μ g of a control cell extract
25 from untreated or healthy cells is loaded on to a SDS-PAGE gel in odd numbered lanes. In adjacent, even numbered lanes, equivalent amounts of experimental extracts are deposited. The latter samples are from cells that have been treated with a hormone or drug or that have been obtained from diseased tissue. The extracts may be prepared by homogenizing cells in buffer containing a detergent such as 0.5% Triton X-100TM and protein phosphatase
30 inhibitors (to preserve the state of protein phosphorylation in the sample). The extracts are then subjected to ultracentrifugation to remove insoluble matter.

To optimize the detection of protein band shifts, the SDS-PAGE gel is precast with a higher than normal concentration of acrylamide and a lower than normal concentration of bisacrylamide. An electric current is applied to the slab gel until proteins with a molecular mass less than 27,000 Dalton are eluted from the bottom of the gel. The proteins remaining on the slab gel are then electro-transferred on to a nitrocellulose or PVDF membrane that traps the proteins. The membrane is cut into separate strips that each contain samples of the resolved proteins from both control and experimental cell extracts. Each strip is probed with a different mixture of primary antibodies (eg. from rabbit) that react with a distinct subset peptide or protein substrate by the protein kinase of interest. Each reaction is conducted in a separate tube or well of a microtitre plate.

One application of this invention is for the discovery of novel protein kinases. The following strategy should permit the rapid acquisition of protein kinase drug targets. The objective of this approach is to identify those protein kinases that demonstrate increased expression or phosphorylation in association with a disease state or in response to an extracellular signal such as mitogen, drug or stress factor. The approach is based on the following:

1. Antibodies developed for one protein kinase can cross-react with structurally related protein kinases.
2. A band shift of a cross-reactive protein on an immunoblot is due to phosphorylation, and increased phosphorylation is probably associated with activation of the kinase. Greater than 90% of the known protein kinases are phosphorylated in their active states. One of the exceptions is glycogen synthase kinase-3, which is inhibited when it is phosphorylated on serine by protein kinase B. However, activation of glycogen synthase kinase-3 is still dependent on tyrosine phosphorylation of this kinase.
3. Proteins that cross-react with protein kinase antibodies and also bind to gamma-ATP-agarose beads have a very high probability of being protein kinases. This resin will capture many ATP binding proteins in addition to protein kinases but this

procedure can purify kinase by up to 200-fold.

4. Proteins that autophosphorylate with [γ -³²P]ATP following immunoprecipitation with protein kinase antibody are likely to be protein kinases. Most antibodies are
5 unsuitable for immunoprecipitation of proteins, and may require partial denaturation of the
proteins. Denatured kinases would have little or no autophosphorylating activity.
5. A combination of gamma-ATP-agarose, immunosorbent and fast protein liquid chromatography column steps followed by SDS-PAGE permits rapid purification of an
10 immunoreactive protein to allow for its identification by sequencing.
6. There is a likelihood that a protein kinase detected with antibodies is novel. Of the
2000 or so kinases expected to exist, only about a quarter have been fully sequenced.
Nevertheless, partial cDNA sequences for most protein kinases are available in public and
15 private EST cDNA sequence databases. Once a portion of the cDNA structure of the
protein kinase gene is available, it is straightforward to obtain the complete nucleotide and
amino acid structures of the gene and its protein with standard methodologies.

One of the beneficial outcomes of this invention is that unknown proteins which can
20 cross-react with the kinase-specific antibodies are detected. Those unidentified proteins that
change in their abundance or their phosphorylation state in response to a disease condition
or treatment are worthy of closer analysis. If such proteins can be shown to bind to
ATP-agarose or capable of autophosphorylation with radioactively labelled ATP, then there
is a high probability that they are protein kinases. Moreover, with the antibody that was
25 originally used to detect a putative kinase, it is possible to rapidly purify the protein so that
it can be sequenced by the Edman degradation method or identified by mass spectroscopy
of trypsin digested fragments of the protein. If any part of the protein has been previously
sequenced, it would be available in public or private protein sequence databases. A partial
sequence in the human EST sequence database may be available. From this information, a
30 full length cDNA sequence for the protein could be rapidly obtained using PCR-based
techniques. This would be worthwhile if the cDNA sequence contained conserved kinase

catalytic subdomain sequences. In this manner, novel protein kinases that display desirable characteristics (eg. increased expression in solid tumour relative to adjacent, normal tissue) can be detected and identified. If the inappropriate activity of such protein kinase is shown to contribute to the development of the disease, then they would be most valuable drug 5 targets.

Measurement of the activation of a protein kinase by this invention is dependent on the detection of its band shift on SDS-PAGE gels. A limited number of protein kinases do not exhibit a band shift change when they are activated. However, their *in vivo* substrates can display band shifts upon their phosphorylation. This can be exploited for the 10 development of *in vitro* and *in vivo* substrate assays. Current approaches for high throughput screening of protein kinase inhibitors *in vitro* involve the use of radioactive [γ -³²]ATP and measurement of the incorporation of the radioactive phosphate into a peptide or protein substrate by protein kinase of interest. Each reaction is conducted in a separate tube or well of a microtitre plate generating high volumes of radioactive garbage.

15 There are many examples of proteins that are highly specific substrates of particular protein kinases; examples include glycogen phosphorylase for phosphorylase kinase, myosin light chain for myosin light chain kinase, eIF2 α for PKR, MARCKS for protein kinase C, Erk1 and Erk2 for Mek1 and Mek2. Antibodies are commercially available for many of these substrates or may be produced as described above. Such antibodies may be 20 used to probe for the phospho-states of the substrates, as revealed by their mobility on immunoblots of SDS-PAGE gels. These substrates would not have to be purified from crude cellular extracts for use in the protein kinase assays. However, since many of the substrates may already exist in phosphorylated forms in cell extracts, it may be necessary to incubate the extracts with active preparations of protein phosphatases, which can be 25 subsequently inactivated with phosphatase inhibitors prior to the kinase assays. With this method, a crude mixture of active protein kinases may be added to the phosphatase-treated cellular extract in a single tube, and the phosphorylation reaction can commence with the inclusion of non-radioactive ATP. Only catalytic amounts of protein kinases will be necessary, so any phosphorylated substrates that contaminate the preparation of protein 30 kinases will be relatively minor compared to the amounts of the substrates in the phosphatase-treated cell extracts. Any kinases that contaminate the phosphatase-treated cell

extracts would not be a concern, since they are actually desirable. It may be necessary to add a kinase preparation after phosphatase treatment of the substrate extracts, because many protein kinases are inhibited when they are dephosphorylated. After a short suitable incubation time, the reactions can be terminated by addition of SDS-PAGE sample buffer.

5 Such substrate analysis can be performed as described above for protein kinases, except that panels of antibodies for the kinase substrates will be used in place of the kinase antibody panels. By this approach, the decreased mobility of the kinase substrates will be evident as band shift on the immunoblots in the absence of kinase inhibitors. The presence of specific protein kinase inhibitors would be revealed by the inhibition of the appearance of the upper
10 bands.

In vitro substrate analysis according to this invention would be ideal for the further characterization of compounds that have already been shown to display inhibitor activity toward a kinase and the selectivity of these compounds is in question. A distinct advantage of this method is that it would be easy to compare the findings with a substrate analysis
15 *in vivo* assay performed using the same blends and concentrations of kinase substrate antibodies that work in the *in vitro* kinase assay. However, the analysis would be performed on extracts from cells that have been incubated with agonists that stimulate the kinases of interest. These cells would also be exposed to the compounds that exhibit inhibitory activity towards kinase *in vitro*. In this manner, the efficacy of these inhibitors
20 could be evaluated in living cells.

By way of illustration, examples from different cell types are given to demonstrate the present invention. The following examples are not intended to be limiting of the invention.

25 **Example 1**

In this example, the kinase multi-blot analysis is used to probe for the presence of over 45 different protein kinases in soluble extracts prepared from the whole brain, heart and skeletal muscle of adult male Sprague-Dawley rats. The results demonstrate
30 large differences in kinase expression patterns between these tissues.

Materials

Affinity-purified rabbit polyclonal or monoclonal antibodies and immunizing peptides used to raise these antibodies are listed in Table 1. These antibodies were either 5 prepared or obtained commercially. Goat anti-rabbit IgG conjugated to alkaline phosphatase (AP) was obtained from Calbiochem (San Diego, CA). Enhanced chemiluminescence (ECL) detection reagents for immunoblotting were obtained from Amersham Pharmacia Biotech, Inc. (Baie d'Urfe, Quebec). Other reagents were obtained from Sigma-Aldrich (St. Louis, MI), unless otherwise stated.

10

Preparation of rat tissue extracts

Brains, hearts and hind leg tibial skeletal muscles from 50-day old male Sprague-Dawley rats were rapidly excised, after induction of anesthesia by intraperitoneal injection of 15 pentobarbital (60 mg/kg). The tissues were cut, rinsed with phosphate buffered saline at 4 °C, frozen in liquid nitrogen, and stored at -70 °C until use. The tissues were pulverized with 5 strokes of a liquid nitrogen-cooled hand French press and re-suspended in 10 volumes of ice-cold homogenization buffer containing: 20 mM MOPS, 15 mM EGTA, 2 mM Na₂EDTA, 1 mM Na₃VO₄, 1 mM dithiothreitol, 75 mM β-glycerophosphate, 0.1 mM 20 phenylmethanesulfonyl fluoride, 1 µg/ml aprotinin, 0.7 µg/ml pepstatin, 1 µg/ml leupeptin, and 1% Triton X-100. This was then sonicated with a Branson Probe Sonicator at 4°C with 3 x 30 s bursts. The homogenates were ultracentrifuged at 100,000 rpm (240,000 x g) for 15 min in a Beckman TLA-100.2 ultracentrifuge at 4 °C. The supernatants were immediately frozen at -70 °C until subsequent analysis.

Table 1. Sources of antibodies.

Kinase	Ab Name	Conc. (μ g/ml)	Source & Catalog No. or immunizing peptide ¹
1. Calmodulin-dependent kinase kinase	CaMKK-CT	0.2	StressGen KAP-CA001
2. Calmodulin-dependent kinase 4	CaMPK4-NT	1.0	StressGen KAP-CA003
3. Cyclin-dependent kinase 1 (cdc2)	Cdc2-CT	0.7	StressGen KAP-CC001
4. Cyclin-dependent kinase 2	Cdk2	1.6	StressGen KAP-CC007E
5. Cyclin-dependent kinase 5	Cdk5-CT	2.4	Upstate 06-258
6. Cyclin-dependent kinase 6	Cdk6	0.8	StressGen KAP-CC006E
7. Cyclin-dependent kinase -7	MO15-PCT	0.5	StressGen KAP-CC010E
8. Cyclin-dependent kinase 8	Cdk8-NT	2.0	StressGen KAP-CC008E
9. Casein kinase 1 α and ϵ	CK1 SG	1.0	StressGen KAP-ST103E
10. Casein kinase 2 α	CK2 α -III	1.2	StressGen KAP-ST010
11. Cot (Tpl2)	Cot-PCT	0.5	EESEMLKRQRSLYIDGC ²
12. p43 and p45 MAP kinase homologues	Erk1-III	0.6	Upstate 06-183
13. Erk1 and Erk2 MAP kinases	Erk1-CT	0.3	Upstate 06-182
14. Erk5 MAP kinase (Bmk)	Erk5-PNT	4.0	SAEPPAREGRTTRPHRC ²
15. Glycogen synthase kinase β	GSK3 β -XI	2.0	StressGen KAP-ST002E
16. Integrin linked kinase 1	ILK1 SG	1.0	StressGen KAP-ST203
17. Kkialre Cdk-like kinase	Kkialre-CT	3.0	StressGen KAP-CC003
18. Kinase-suppressor of Ras	Ksr1-CT	1.0	EKLPKLNRRRLSHPGHFWKSC ²
19. MAP kinase-activated kinase 2	MAPKAPK2-PCT	1.0	StressGen KAP-MA015E
20. MAP kinase kinase 1	Mek1-XI	2.0	Upstate 06-235
21. MAP kinase kinase 3	Mkk3-CT	1.0	Upstate 06-615
22. MAP kinase kinase 4	Mkk4-XI	1.6	Upstate 06-281
23. MAP kinase kinase 5	Mek5-PNT	1.6	StressGen KAP-MA003
24. MAP kinase kinase 6	Mek6-SG	0.8	StressGen KAP-MA014E
25. MAP kinase kinase kinase 1	Mekk1-PNT	1.2	StressGen KAP-SA010
26. MAP kinase kinase kinase 3	Mekk3 SG	0.5	StressGen KAP-MA013E
27. Mos	Mos-III	1.0	StressGen KAP-MA004
28. p38 MAP kinase	p38 Hog-CT	0.2	StressGen KAP-MA009E
29. p21-activated kinase α	Pak (C-19)	0.8	SantaCruz sc-881
30. Pim1	Pim1-T	1.4	StressGen KAP-ST004
31. Protein kinase A (cAMP-dep. kinase)	PKA-NT	3.0	StressGen KAP-PK001
32. Protein kinase C β	PKC- β M7 mAb	0.5	Gift from Susan Jaken ³
33. Protein kinase C ϵ	nPKC- ϵ (C-15)	0.5	SantaCruz sc-214
34. Protein kinase C ζ	PKC- ζ (C-20)	0.5	SantaCruz sc-226
35. Protein kinase G (cGMP-dep. kinase)	PKG1-CT	2.6	StressGen KAP-PK005
36. Protein kinase B α	PKB-CT	1.8	Upstate 06-276
37. Protein kinase B β	PKB2-PCT	2.0	CRYDSLGLLEDQRT ²
38. RafB	RafB-CT	3.0	StressGen KAP-MA006
39. Ribosomal S6 kinase 1	Rsk1 (C-21)	0.7	SantaCruz sc-231
40. Ribosomal S6 kinase 2	Rsk2-PCT	0.8	StressGen KAP-ST007
41. S6 protein kinase	S6K-PNT	1.8	Upstate 06-321
42. Stress-activated kinase (Jnk)	SAPK β	1.6	StressGen KAP-SA004
43. TGF β -activated kinase	Tak1-CT	1.0	StressGen KAP-ST009E

Notes

(1) Upstate = Upstate Biotechnology Inc. (Lake Placid, New York, U.S.A.); StressGen = StressGen Biotechnologies Corp. (Victoria, British Columbia, Canada); Santa Cruz Biotechnology, Inc. (Santa Cruz, California, U.S.A.).

(2) All of these antibodies are commercially available, except for Cot-PCT, Erk5-PNT, Ksr1-CT, PKC- β M7 and PKB2-PCT. For these antibodies, the amino acid sequence of the immunizing peptide is provided.

(3) All of the antibodies used were rabbit polyclonal antibodies with the exception of the PKC- β M7 mouse monoclonal antibody provided by Dr. Susan Jaken's of the W. Alton Jones Cell Science Center in Lake Placid, New York.

Gel electrophoresis and immunoblotting

The thawed cell lysates were measured for protein content using Bradford reagent (Bio-Rad) with bovine serum albumin as the reference standard. The protein concentration of the lysates was adjusted to 1 mg/ml in SDS-PAGE sample buffer (2% SDS, 5% glycerol, 50 mM Tris-HCl, pH 6.8, 0.1 M β -mercaptoethanol and 0.01% bromophenol blue) and boiled at 100 °C for 3 min. One mg of the cell lysate was loaded on to the stacking layer (2 mm x 4 cm x 20 cm; 4% acrylamide/0.11% bisacrylamide) of an SDS-PAGE gel. A comb was not used to create individual lanes, so that there was 10 a single, wide lane over the width of the entire gel. The stacking gel was previously layered over a separating SDS-PAGE gel (2 mm x 12.5 cm x 20 cm; 13% acrylamide/0.086% bisacrylamide). Electrophoresis was performed at 30 mA (maximum voltage) and was continued until proteins of 25,000 Daltons had migrated to the bottom of the gel. The composition and concentrations of the other ingredients in the 15 stacking and separate gels, and in the lower and upper chamber gel buffers were as described (Laemmli, U.K. (1970) *Nature* 227, 680-685). Proteins were then electrophoretically transferred from the gel at 300 mA (maximum voltage) for 3 h on to a nitrocellulose membrane, and the membrane was subsequently cut vertically into 1 cm wide strips. The strips were then blocked with 5% skim milk powder in Tris-buffered 20 saline (20 mM Tris/HCl, pH 7.5, 0.5 M NaCl, 0.2 % Tween-20; TBST) and, after quickly rinsing the membrane with TBST, each strip was exposed to a unique mixture of different primary antibodies in TBST for 3 h with constant shaking at room temperature. The concentrations of the antibodies that were used are provided in Table 1. The strips were washed two times for 15 min with TBST and incubated with horse 25 radish peroxidase-conjugated secondary antibody (goat anti-rabbit or anti-mouse IgG) in TBST for 30 min. After washing the strips three times for 10 min with TBST, the strips were reassembled, and subjected to the ECL Western blotting detection system (Amersham Pharmacia Biotech, Inc.). Exposure of the x-ray films was for 40 sec.

Results

Figure 1 shows an example of the application of multi-kinase immunoblotting technique applied to three different rat tissues. It is evident that the patterns of kinase expressions differed markedly between the tissues. At least 45 known protein kinases were visualized on the immunoblots and clearly identified based on their predicted sizes and immunoreactivities. From the intensity of the signals for the immunoreactive kinase bands in Fig. 1, the relative expression levels of these specific protein kinases are provided in Table 2. This sort of analysis could be especially useful for tracking kinases as a function of development, long term mitogen, stress or drug stimulation, and disease progression.

Table 2. Relative expression of known protein kinases in rat tissues.

Kinase	Relative Expression	Brain	Heart	Skeletal muscle
1. Calmodulin-dependent kinase kinase	Moderate	Low	Low	
2. Calmodulin-dependent kinase 4	Moderate	None	None	
3. Cyclin-dependent kinase 1 (cdc2)	Low	None	None	
4. Cyclin-dependent kinase 2	None	Low	None	
5. Cyclin-dependent kinase 5	Moderate	None	None	
6. Cyclin-dependent kinase 6	Low	Moderate		Moderate
7. Cyclin-dependent kinase 7	Moderate	Low		Low
8. Cyclin-dependent kinase 8	Low	Low		Low
9. Casein kinase 1 α	Low	Low		Low
10. Casein kinase 1 ϵ	Moderate	Low		Low
11. Casein kinase 2 α	High	High		High
12. Cot (Tp12)	Low	High		Low
13. Erk1 MAP kinase	High	High		High
14. Erk2 MAP kinase	High	High		High
15. Erk5 MAP kinase (Bmk)	Moderate	None		None
16. Glycogen synthase kinase β	High	High		Moderate
17. Integrin linked kinase 1	Moderate	Moderate		Moderate
18. Kkialte Cdk-like kinase	None	Moderate		High
19. Kinase-suppressor of Ras	Low	None		None
20. MAP kinase-activated kinase 2	High	Low		None
21. MAP kinase kinase 1 (Mek1)	High	Low		Low
22. MAP kinase kinase 3 (Mek3)	Low	Low		Low
23. MAP kinase kinase 4 (Mek4)	High	Low		Low
24. MAP kinase kinase 5 (Mek5)	High	Moderate		Low
25. MAP kinase kinase 6 (Mek6)	High	High		High
26. MAP kinase kinase kinase 1 (Mekk1)	Low	Moderate		Low
27. MAP kinase kinase kinase 3 (Mekk3)	Low	Low		Low
28. Mos	None	Low		Moderate
29. p38 α MAP kinase	Moderate	High		Moderate
30. p21-activated kinase α	High	Low		Low
31. Pim1	Moderate	High		Low
32. Protein kinase A (cAMP-dep. kinase)	Moderate	High		High
33. Protein kinase B α (Akt1)	High	Low		Low
34. Protein kinase B β (Akt2)	High	Low		Low
35. Protein kinase C β	High	Low		Low
36. Protein kinase C ϵ	High	High		High
37. Protein kinase C ζ	High	High		High
38. Protein kinase G (cGMP-dep. kinase)	Low	High		High
39. RafB	High	Low		None
40. Ribosomal S6 kinase 1	High	High		Moderate
41. Ribosomal S6 kinase 2	Low	Low		High
42. S6 protein kinase	High	Low		High
43. Stress-activated kinase β p46 (Jnk)	High	Low		Low
44. Stress-activated kinase β p54 (Jnk)	High	Moderate		moderate
45. TGF β -activated kinase	Low	Low		Low

Example 2

In this example, the kinase multi-blot analysis is used to probe for the presence and 5 activation states of over 45 different protein kinases in soluble extracts prepared from the human Ramos B cell line that have been treated anti-IgM antibody in order to stimulate these cells through a B cell antigen receptor. The results demonstrate the band shifting of several protein kinases as a consequence of their increased phosphorylation in response to B cell antigen receptor stimulation.

10

Materials

Affinity-purified rabbit polyclonal or monoclonal antibodies and immunizing peptides used to raise these antibodies are listed in Table 1. Goat anti-rabbit IgG 15 conjugated to alkaline phosphatase (AP) was obtained from Calbiochem. Enhanced chemiluminescence (ECL) detection reagents for immunoblotting were obtained from Amersham Pharmacia Biotech, Inc. Other reagents were obtained from Sigma-Aldrich, unless otherwise stated.

20 ***Preparation of cell extracts***

The human Ramos B cell line (American Type Culture Collection, Rockville, MD) was cultured in Dulbecco's modified Eagle medium containing 10% heat inactivated fetal bovine serum and 2 mM glutamine at 37°C in a 5% CO₂/air mixture. 25 For each experimental analysis, 2 x 10⁷ cells were seeded in a 150 mm culture dish containing 20 ml of medium. Twelve hours prior to cell stimulation, the cells were cultured in the above media in the absence of serum, and then were incubated for 5 min with anti-IgM antibody. Subsequently the cells were lysed in 2 ml of ice-cold buffer that contained 20 mM MOPS, pH 7.2, 5 mM EGTA, 1% (w/v) Nonidet P-40, 1 mM 30 dithiothreitol, 75 mM β -glycerol phosphate, 1 mM Na₃VO₄, and 1 mM phenylmethylsulfonyl fluoride by sonication with a Branson Probe Sonicator at 4°C with

3 x 30 s bursts. The homogenates were ultracentrifuged at 100,000 rpm (240,000 x g) for 15 min in a Beckman TLA-100.2 ultracentrifuge at 4 °C. The supernatants were immediately frozen at -70 °C until subsequent analysis.

5 *Gel electrophoresis and immunoblotting*

The thawed cell lysates were measured for protein content using Bradford reagent (Bio-Rad) with bovine serum albumin as the reference standard. The protein concentration of the lysates was adjusted to 1 mg/ml in SDS-PAGE sample buffer (2% SDS, 5% glycerol, 50 mM Tris-HCl, pH 6.8, 0.1 M β -mercaptoethanol and 0.01% bromophenol blue) and boiled at 100 °C for 3 min. One mg of the cell lysate was loaded on to the stacking layer (2 mm x 4 cm x 20 cm; 4% acrylamide/0.11% bisacrylamide) of an SDS-PAGE gel. A twenty lane comb was used, and the extracts from untreated (control) and anti-IgM-treated (experimental) cells were deposited into adjacent lanes.

10 Molecular mass markers (glycogen phosphorylase, bovine serum albumin, ovalbumin, glyceraldehyde 3-phosphate dehydrogenase) were applied to the first and last lanes of the gel. The stacking gel was previously layered over a separating SDS-PAGE gel (2 mm x 12.5 cm x 20 cm; 13% acrylamide/0.086% bisacrylamide). Electrophoresis was performed at 30 mA (maximum voltage) and was continued until proteins of 25,000

15 Daltons had migrated to the bottom of the gel. The composition and concentrations of the other ingredients in the stacking and separate gels, and in the lower and upper chamber gel buffers were as described (Laemmli, U.K. (1970) *Nature* 227, 680-685). Proteins were then electrophoretically transferred from the gel at 300 mA (maximum voltage) for 3 h on to a nitrocellulose membrane, and the membrane was subsequently

20 cut vertically into strips that contained one lane of the control and one lane of the experimental samples. The strips were then blocked with 5% skim milk powder in Tris-buffered saline (20 mM Tris/HCl, pH 7.5, 0.5 M NaCl, 0.2 % Tween-20; TBST) and, after quickly rinsing the membrane with TBST, each strip was exposed to a unique mixture of different primary antibodies in TBST for 3 h with constant shaking at room

25 temperature. The concentrations of the antibodies that were used are provided in Table

30 1. The strips were washed two times for 15 min with TBST and incubated with horse

radish peroxidase-conjugated secondary antibody (goat anti-rabbit or anti-mouse IgG) in TBST for 30 min. After washing the strips three times for 10 min with TBST, the strips were reassembled, and subjected to the ECL Western blotting detection system (Amersham Pharmacia Biotech, Inc.). Exposure of the x-ray films was for 40 sec.

5

Results

In the human Ramos B cell line, many of the tested protein kinases were not detected at measurable levels by the kinase multi-blot analysis. Several of the protein kinases present in

10 the Ramos cells were observed to undergo band shifts to high apparent molecular mass species following short term exposure for 5 min to the anti-IgM antibody. These kinases are evident in Fig. 2. This band shifting generally correlated with the phosphorylation and activation of these kinases. Table 3 summarizes the findings from this experiment with respect to the expression level and band shifting of the 45 known protein kinases that were examined.

15 An unidentified 52-kDa immunoreactive protein (p52) in Lanes 1 and 2 that band shifted upon anti-IgM antibody treatment may correspond to a novel protein kinase within the MAP kinase family based on immunoreactivity.

Table 3. Protein kinase expression and band shifting in anti-IgM antibody-treated Ramos cells.

Kinase	Expression Level	Band Shifted
1. Calmodulin-dependent kinase kinase	None	
2. Calmodulin-dependent kinase 4	None	
3. Cyclin-dependent kinase 1 (cdc2)	None	
4. Cyclin-dependent kinase 2	None	
5. Cyclin-dependent kinase 5	None	
6. Cyclin-dependent kinase 6	None	
7. Cyclin-dependent kinase 7	None	
8. Cyclin-dependent kinase 8	None	
9. Casein kinase 1 α	High	No
10. Casein kinase 1 ϵ	None	
11. Casein kinase 2 α	High	No
12. Cot (Tpl2)	High	No
13. Erk1 MAP kinase	High	Yes
14. Erk2 MAP kinase	High	?
15. Erk5 MAP kinase (Bmk)	None	
16. Glycogen synthase kinase β	None	
17. Integrin linked kinase 1	None	
18. Kkialre Cdk-like kinase	None	
19. Kinase-suppressor of Ras	None	
20. MAP kinase-activated kinase 2	High	No
21. MAP kinase kinase 1 (Mek1)	High	No
22. MAP kinase kinase 3 (Mek3)	None	
23. MAP kinase kinase 4 (Mek4)	None	
24. MAP kinase kinase 5 (Mek5)	Low	No
25. MAP kinase kinase 6 (Mek6)	Moderate	No
26. MAP kinase kinase kinase 1 (Mekk1)	None	
27. MAP kinase kinase kinase 3 (Mekk3)	None	
28. Mos	None	
29. p38 α MAP kinase	Moderate	No
30. p21-activated kinase α	None	
31. Pim1	None	
32. Protein kinase A (cAMP-dep. kinase)	None	
33. Protein kinase B α (Akt1)	None	
34. Protein kinase B β (Akt2)	High	Yes
35. Protein kinase C β	High	Yes
36. Protein kinase C ϵ	High	Yes
37. Protein kinase C ζ	None	
38. Protein kinase G (cGMP-dep. kinase)	None	
39. RafB	None	
40. Ribosomal S6 kinase 1	High	Yes
41. Ribosomal S6 kinase 2	None	
42. S6 protein kinase	Moderate	Yes
43. Stress-activated kinase β p46 (Jnk)	None	
44. Stress-activated kinase β p54 (Jnk)	Moderate	No
45. TGF β -activated kinase	High	No

Example 3

In this example, the kinase multi-blot analysis is used to probe for the effect of various protein kinase inhibitors on the ability of hepatocyte growth factor (HGF) to 5 activate protein kinases in the Erk1/Erk2, p38 MAP kinase and S6 kinase pathways. This example demonstrates that unique kinase band shift patterns may be produced by different drugs. This could be exploited to determine the mechanisms of action of known drugs and the identification of unknown targets of new drugs.

10 *Materials*

The protein kinase inhibitors PD98059 (for Mek1), LY294002 (for phosphatidylinositol 3-kinase), SB203580 (for p38 MAP kinase) and rapamycin (for mTor/FRAP) were obtained from ProMega Corporation (Madison, WI). Affinity-15 purified rabbit polyclonal or monoclonal antibodies and immunizing peptides used to raise these antibodies are listed in Table 1. Goat anti-rabbit IgG conjugated to alkaline phosphatase (AP) was obtained from Calbiochem. Enhanced chemiluminescence (ECL) detection reagents for immunoblotting were obtained from Amersham-Pharmacia Biotech, Inc. Other reagents were obtained from Sigma-Aldrich, unless otherwise 20 stated.

Preparation of cell extracts

Primary cultures of human ovarian surface epithelial cells were cultured in 25 Dulbecco's modified Eagle medium containing 10% heat inactivated fetal bovine serum and 2 mM glutamine at 37°C in a 5% CO₂/air mixture. For each experimental analysis, 2 x 10⁷ cells were seeded in a 150 mm culture dish containing 20 ml of medium. Twelve hours prior to cell stimulation, the cells were cultured in the above media in the absence of serum, and then were incubated at 37°C in the absence or presence of either 50 µM 30 PD98059, 50 µM LY294002, 10 µM SB203580 or 20 nM rapamycin for 30 min. The cells were subsequently incubated for another 10 min with 20 ng/ml of HGF prior to

their lysis at 4°C in 2 ml of buffer that contained 20 mM MOPS, pH 7.2, 5 mM EGTA, 1% (w/v) Nonidet P-40, 1 mM dithiothreitol, 75 mM β -glycerol phosphate, 1 mM Na₃VO₄, and 1 mM phenylmethylsulfonyl fluoride. The cells were sonicated with a Branson Probe Sonicator at 4°C with 3 x 30 s bursts. The homogenates were 5 ultracentrifuged at 100,000 rpm (240,000 x g) for 15 min in a Beckman TLA-100.2 ultracentrifuge at 4 °C. The supernatants were immediately frozen at -70 °C until subsequent analysis.

Gel electrophoresis and immunoblotting

10

The thawed cell lysates were measured for protein content using Bradford reagent (Bio-Rad) with bovine serum albumin as the reference standard. The protein concentration of the lysates was adjusted to 1 mg/ml in SDS-PAGE sample buffer (2% SDS, 5% glycerol, 50 mM Tris-HCl, pH 6.8, 0.1 M β -mercaptoethanol and 0.01% 15 bromophenol blue) and boiled at 100 °C for 3 min. One mg of the cell lysate was loaded on to the stacking layer (2 mm x 4 cm x 20 cm; 4% acrylamide/0.11% bisacrylamide) of an SDS-PAGE gel. A twenty lane comb was used, and the extracts from untreated (control; Lane 1), HGF-treated (Lane 2), PD98059- and HGF-treated (Lane 3), LY294002- and HGF-treated (Lane 4), SB203580- and HGF-treated (Lane 5), and 20 rapamycin- and HGF-treated (Lane 6) cells were deposited into adjacent lanes. Molecular mass markers (glycogen phosphorylase, bovine serum albumin, ovalbumin, glyceraldehyde 3-phosphate dehydrogenase, trypsinogen) were applied to the first and last lanes of the gel. The stacking gel was previously layered over a separating SDS-PAGE gel (2 mm x 12.5 cm x 20 cm; 13% acrylamide/0.086% bisacrylamide). 25 Electrophoresis was performed at 30 mA (maximum voltage) and was continued until proteins of 25,000 Daltons had migrated to the bottom of the gel. The composition and concentrations of the other ingredients in the stacking and separate gels, and in the lower and upper chamber gel buffers were as described (Laemmli, U.K. (1970) *Nature* 227, 680-685). Proteins were then electrophoretically transferred from the gel at 300 30 mA (maximum voltage) for 3 h on to a nitrocellulose membrane, and the membrane was subsequently cut vertically into strips that contained six lanes of the control and various

HGF and inhibitor-treated cells. The strips were then blocked with 5% skim milk powder in Tris-buffered saline (20 mM Tris/HCl, pH 7.5, 0.5 M NaCl, 0.2 % Tween-20; TBST) and, after quickly rinsing the membrane with TBST, each strip was exposed to a unique mixture of different primary antibodies in TBST for 3 h with constant shaking at room temperature. The concentrations of the antibodies that were used are provided in Table 1. The strips were washed two times for 15 min with TBST and incubated with horse radish peroxidase-conjugated secondary antibody (goat anti-rabbit or anti-mouse IgG) in TBST for 30 min. After washing the strips three times for 10 min with TBST, they were subjected to the ECL Western blotting detection system (Amersham Pharmacia Biotech, Inc.). Exposure of the x-ray films was for 40 sec.

Results

Many different protein kinases were clearly detected in the human ovarian surface epithelial (OSE) cells. For the purposes of this example, only the band shifted states of six protein kinases are presented in Figure 3. These kinases are the extracellular regulated kinases Erk1 (Panel A) and Erk2 (Panel A) isoforms, p38 Hog MAP kinase (Panel B), protein kinase B alpha (PKB1, Panel C) and beta (PKB2, Panel D) isoforms, and p70 S6 kinase (S6K). A schematic of the signalling pathways within which these kinases participate is present on the right in Figure 3. On the one hand, HGF is known to stimulate the enzyme activities of the Erk1 and Erk2 MAP kinases as well as p70 S6K. On the other hand, there are no reports of activation of p38 MAP kinase by HGF. In concordance, HGF treatment of the OSE cells produced reduced mobilities of Erk1, PKB1, PKB2 and S6K, consistent with their phosphorylation and activation. A band shift in Erk2 was not evident due to comigration of activated Erk2 with the inactive form of Erk1. There was no change in the mobility in p38 MAP kinase. Consistent with the known action of PD98059, the Mek1 inhibitor prevented the HGF-induced band shift in Erk1, but not of PKB1 and PKB2. There was some reduction of the HGF-induced S6K band shift, possibly because some of the phosphorylation of S6K may be catalyzed by Erk1 and Erk2. The phosphatidylinositol 3-kinase (PI3K) inhibitor LY294002 caused a slight reduction in HGF-induced Erk1

band shifting, in part because the protein kinase C zeta (PKC ζ) isoform is normally activated by the lipid products of the PI3K reaction, and PKC ζ causes the activation of Mek1. The inhibition of PI3K by LY294002 completely blocked the HGF-induced band shifting of PKB1, PK2 and S6K, consistent with their actions distal to PI3K.

5 Furthermore, the LY294002 produced band shifts in these kinases to forms of lower molecular mass than were detected in the untreated cells. These findings indicated that there was an intermediate state of activation of these kinases in the control cells in the absence of HGF. The p38 MAP kinase inhibitor SB203580 had no discernable effect on any of the kinases in the absence or presence of HGF. The mTOR/Frap inhibitor

10 rapamycin only reduced the band shift in S6K, which is expected since this protein kinase appears to lie upstream of only the S6K and none of the other protein kinase that were tracked.

One application of the multi-kinase analysis is for drug profiling. It is evident in

15 Example 3, that the different protein kinase inhibitors generated distinct changes in the 6 protein kinases that were tracked. Potentially several hundred protein kinases can be monitored for the specific effects of selected drugs. The short term actions of the drug on the basal, mitogen-stimulated and stress-stimulated phosphorylation states of the various protein kinases can be assessed. Distinct sets of band shifts should be produced by different drugs.

20 These patterns can be interpreted to deduce the mechanisms of action of these drugs. By matching the patterns of kinase alterations induced by known drugs, it would be possible to determine targets of unknown drugs. If two drugs generate exactly the same patterns of kinase changes, then they should have the same cellular target.

25 *Example 4*

In this example, the kinase multi-blot analysis is used to probe for changes in the expression of protein kinases in extracts from human breast tumours compared to patient-matched "normal" breast tissue. The method allows for the detection of known

30 protein kinases and other proteins that are increased in samples from diseased tissues or cells. These proteins can serve as markers of disease progression and possibly targets

for therapeutic intervention.

Materials

5 Affinity-purified rabbit polyclonal or monoclonal antibodies and immunizing peptides used to raise these antibodies are listed in Table 1. Goat anti-rabbit IgG conjugated to alkaline phosphatase (AP) was obtained from Calbiochem. Enhanced chemiluminescence (ECL) detection reagents for immunoblotting were obtained from Amersham Pharmacia Biotech, Inc. Other reagents were from Sigma-Aldrich, unless
10 otherwise stated.

Tissue procurement and homogenization

Breast tumours from patients and their adjacent control samples were obtained
15 through the Pathology Department at Vancouver General Hospital. The samples were immediately placed in liquid nitrogen until analysis. Homogenization was carried out by placing the tissue in 5 ml of buffer containing 20 mM MOPS, 50 mM β -glycerophosphate, 1% NP40, 50 mM sodium fluoride, 1 mM Na_3VO_4 , 5 mM EGTA, 2 mM EDTA, 1 mM dithiothreitol, 1 mM benzamidine and 1 mM
20 phenylmethanesulphonyl fluoride, 10 g/ml of leupeptin and aprotinin, and applying three, 20 second bursts of a Brinkman Polytron at a setting of 10,000 rpm whilst on ice. The samples were then subjected to centrifugation at 150,000 x g for 30 min and the supernatant fractions were stored at -70°C until used.

25 Gel electrophoresis and immunoblotting

The thawed cell lysates were measured for protein content using Bradford reagent (Bio-Rad) with bovine serum albumin as the reference standard. The protein concentration of the lysates was adjusted to 1 mg/ml in SDS-PAGE sample buffer (2%
30 SDS, 5% glycerol, 50 mM Tris-HCl, pH 6.8, 0.1 M -mercaptoethanol and 0.01% bromophenol blue) and boiled at 100 °C for 3 min. One mg of the cell lysate was loaded

on to the stacking layer (2 mm x 4 cm x 20 cm; 4% acrylamide/0.11% bisacrylamide) of an SDS-PAGE gel. A twenty lane comb was used, and the extracts from patient-matched normal (control) and tumour breast tissue biopsy extracts were deposited into adjacent lanes. The stacking gel was previously layered over a separating SDS-PAGE 5 gel (2 mm x 12.5 cm x 20 cm; 13% acrylamide/0.086% bisacrylamide). Electrophoresis was performed at 30 mA (maximum voltage) and was continued until proteins of 25,000 Daltons had migrated to the bottom of the gel. The composition and concentrations of the other ingredients in the stacking and separate gels, and in the lower and upper chamber gel buffers were as described (Laemmli, U.K. (1970) *Nature* 227, 680-685).
10 Proteins were then electrophoretically transferred from the gel at 300 mA (maximum voltage) for 3 h on to a nitrocellulose membrane, and the membrane was subsequently cut vertically into strips that contained one lane of the control and one lane of the experimental samples. The strips were then blocked with 5% skim milk powder in Tris-buffered saline (20 mM Tris/HCl, pH 7.5, 0.5 M NaCl, 0.2 % Tween-20; TBST) and, 15 after quickly rinsing the membrane with TBST, each strip was exposed to a unique mixture of different primary antibodies in TBST for 3 h with constant shaking at room temperature. The concentrations of the antibodies that were used are provided in Table 1. The strips were washed two times for 15 min with TBST and incubated with horse radish peroxidase-conjugated secondary antibody (goat anti-rabbit or anti-mouse IgG) in 20 TBST for 30 min. After washing the strips three times for 10 min with TBST, they were subjected to the ECL Western blotting detection system (Amersham Pharmacia Biotech, Inc.). Exposure of the x-ray films was for 2 min.

Results

25 Many different known protein kinases were clearly detected in the human breast tumours and patient matched "normal" breast biopsy samples. Most of these kinases were unaltered in their relative expression levels. Five protein kinases that consistently exhibited increased amounts in the tumour samples are presented in set of left panels in 30 Figure 4. These kinases were p38 Hog MAP kinase, protein kinase B-alpha (PKB α), the various isoforms of the catalytic subunit of casein kinase 2 (CK2), cGMP-dependent

protein kinase (PKG) and cyclin-dependent kinase 8 (Cd8). The elevated amounts of one or more of these kinases may contribute to the neoplastic transformation of breast tissue. Alternatively, they may represent feedback responses to counteract the loss of growth control in the breast tumours. In either event, they may serve as useful markers of
5 cancer progression.

Many antibodies are able to cross-react with related proteins that share the epitopes that are recognized for binding by these antibodies. Since most protein kinases are evolutionarily related, there is a high probability of cross-reactivity of kinase-directed
10 antibodies with homologous kinases. In the experiment described in this example, there were at least 12 unidentified immunoreactive proteins with kinase antibodies that were selectively detected in the tumour samples, but poorly if at all the control samples. Five of these proteins are shown in the immunoblots in the set of panels on the right side of Figure 4. It is possible to identify these proteins following their enrichment by standard purification techniques and
15 protein microsequence analysis. Three major advantages of kinase purification with antibodies are: (1) it is unnecessary to know what will serve as a selective substrate to monitor the presence of the kinase during its purification; (2) it is not critical to preserve the enzyme activity of the kinase during the purification procedures; and (3) the position of the kinase can be clearly detected on an SDS-PAGE gel by immunoblotting for its sequencing, even if the
20 protein is not completely pure.

Example 5

In this example, standard immunoblotting of 2D isoelectric focusing/SDS-PAGE
25 gels of rat brain extracts was undertaken to demonstrate the limitations of the traditional approach for proteomic analysis as applied to protein kinases.

Materials

30 The Immobiline DryStrip Kit with immobilized IPG ampholytes was obtained from Amersham Pharmacia Biotech, Inc. The rabbit polyclonal antibodies Erk1-CT

(Catalogue No. 06-182) and PKC-III (Catalogue No. KAP-PK003) were obtained from Upstate Biotechnology Inc. and StressGen Biotechnologies Corp., respectively. Goat anti-rabbit IgG conjugated to alkaline phosphatase (AP) were obtained from Calbiochem. Enhanced chemiluminescence (ECL) detection reagents for 5 immunoblotting were obtained Amersham Pharmacia Biotech, Inc. Other reagents were from Sigma-Aldrich, unless otherwise stated.

Preparation of rat brain extract

10 The brain from a 50-day old male Sprague-Dawley rat was rapidly excised, after induction of anesthesia by intraperitoneal injection of pentobarbital (60 mg/kg). The brain was cut, rinsed with phosphate buffered saline at 4 °C, and re-suspended in 10 volumes of ice-cold homogenization buffer containing: 10 mM Tris, pH 7.2, 0.1 mM EGTA, 0.1 mM Na₂EDTA, 1 mM Na₃VO₄, 50 mM dithiothreitol, 10 mM β -
15 glycerophosphate, 1 mM phenylmethanesulfonyl fluoride, 10% glycerol and 1% Triton X-100. This was then homogenized with a Brinkman Polytron homogenizer. The homogenates were ultracentrifuged at 100,000 rpm (240,000 x g) for 15 min in a Beckman TLA-100.2 ultracentrifuge at 4 °C. The supernatant was immediately frozen at -70 °C until subsequent analysis.

20

Gel electrophoresis and immunoblotting

25 The thawed cell lysate was measured for protein content using Bradford reagent (Bio-Rad) with bovine serum albumin as the reference standard. One mg of the soluble protein was applied to an Immobiline isoelectric focusing (IEF) DryStrip and electrophoresis was performed on Hoefer SE600 standard vertical gel system as recommended by the manufacturer (Pharmacia Biotech). The 18 cm, pH 3-10 DryStrip was previously rehydrated in 6 M urea, 2 M thiourea, 4% CHAPS, 0.02% bromophenol blue, 2% Nonidet P-40, 0.7% dithiothreitol, 10 mM Tris, 2% IPG 30 ampholyte, 10% glycerol, and 4 mM tributyl phosphine. Following the isoelectric focusing, the IPG strip was overlaid lengthwise on to an 11% SDS-polyacrylamide gel,

and electrophoresis was continued into the second dimension. In a separate lane at the left side of the same 11% SDS-PAGE gel, 200 g of the cytosolic protein was directly loaded on to the gel prior to the electrophoresis into the second dimension. Subsequently, the resolved proteins were then electrophoretically transferred from the 5 2D gel at 300 mA (maximum voltage) for 3 h on to a nitrocellulose membrane. The membrane was then blocked with 5% skim milk powder in Tris-buffered saline (20 mM Tris/HCl, pH 7.5, 0.5 M NaCl, 0.2 % Tween-20; TBST) and, after quickly rinsing the membrane with TBST, it was exposed to a mixture of 0.2 g/ml of Erk1-CT and 1 mg/ml of the PKC-III primary antibodies in TBST for 3 h with constant shaking at room 10 temperature. The membrane was washed two times for 15 min with TBST and incubated with horse radish peroxidase-conjugated secondary antibody (goat anti-rabbit IgG) in TBST for 30 min. After washing the membrane three times for 10 min with TBST, it was subjected to the ECL Western blotting detection system (Amersham Pharmacia Biotech, Inc.). Exposure of the x-ray film was for 5 min.

15

Results

The migration positions of PKC, Erk1 and Erk2 are shown in Figure 5. The right most of the series of spots for Erk1 and Erk2 correspond to their fully 20 dephosphorylated states, and these proteins shifted progressively to the left with the acquisition of each phosphate group. Only one large spot was evident for PKC β . The large black smear on the left of the 2D gel corresponds to five-fold less brain extract applied directly to the same SDS-PAGE gel than was also loaded on to the IPG Drystrip, which was then transferred into the SDS-PAGE gel. These results were 25 consistently observed in multiple experiments, even when diverse agents were used to try to release the proteins from the IPG Drystrip. The adsorption of protein to the IPG Drystrip was especially problematic for less abundant cellular proteins such as protein kinases. Evidently, less than 5% of the Erk1 and Erk2 that loaded on to the IPG Drystrip actually entered the SDS-PAGE gel when electrophoresis was performed in the 30 second dimension.

The unpredictable retention of protein kinases and their substrates by the IPG Drystrips calls into question the useful of standard 2D gel electrophoresis (eg. according to O'Farrell) for resolution of proteins to perform quantitative studies to monitor 5 changes in protein expression and post-translational covalent modification. It is possible that the phosphorylation state of proteins may also influence their binding to the IPG Drystrips. This would compromise on the analysis of the relative amounts of the dephosphorylated to phosphorylated species of some proteins on 2D gels. It is possible that the problems associated with protein retention on the IPG Drystrips with 10 immobilized ampholytes might be avoided through the use of tube gels and soluble ampholytes. However, the protein loading capacity of such tube gels is markedly lower. This could still result in insufficient protein on the 2D gel for detection of protein kinases by silver staining or immunostaining.

An attempt was made to detect more than twenty different protein kinases by 15 immunoblotting 2D gels of rat brain lysates with the antibodies described in Table 1. The immunoblots were performed with mixes composed of two to four different antibodies. With few exceptions, it was very difficult to detect any spots that corresponded to the appropriate sizes of the target proteins. The data shown in Figure 5 represents the best finding using particularly sensitive antibodies. 2D gel electrophoresis is thus insufficiently sensitive and 20 inconsistently reproducible for the simultaneous detection of rare proteins such as protein kinases.

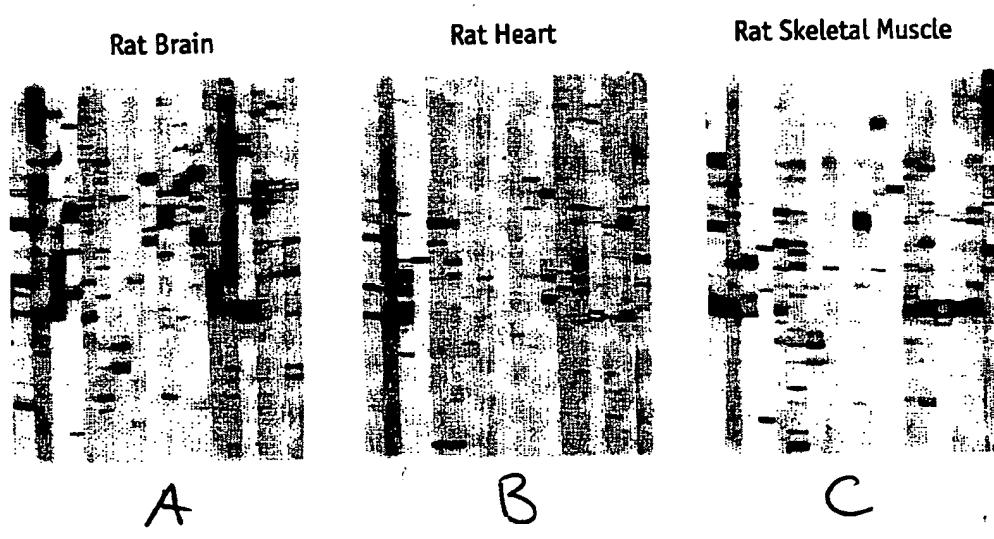


FIG. 1

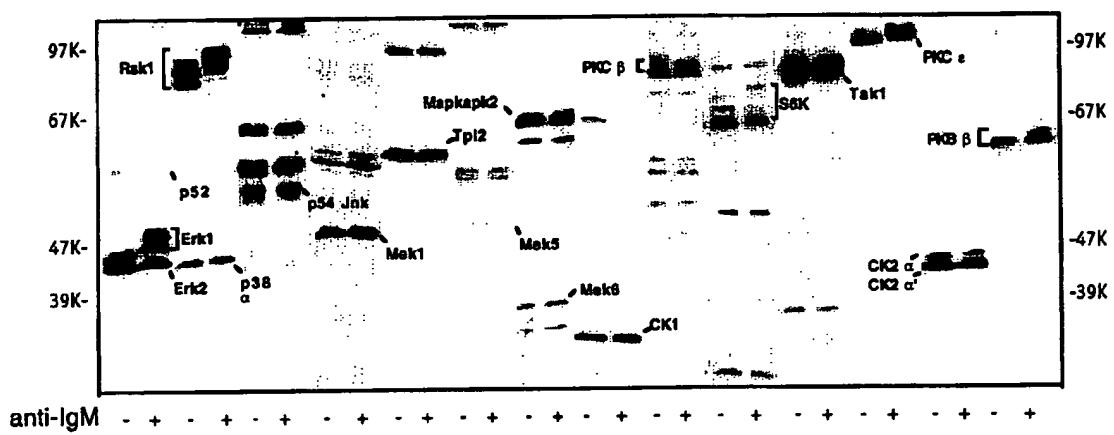


FIG. 2

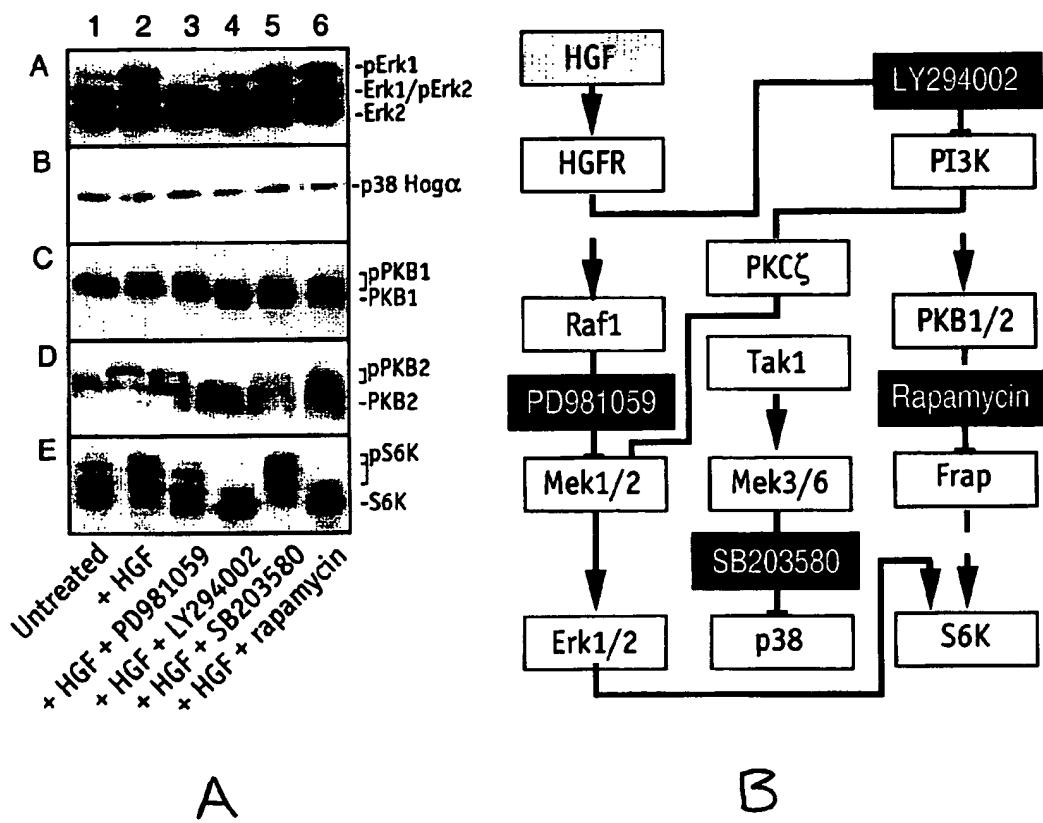


FIG. 3

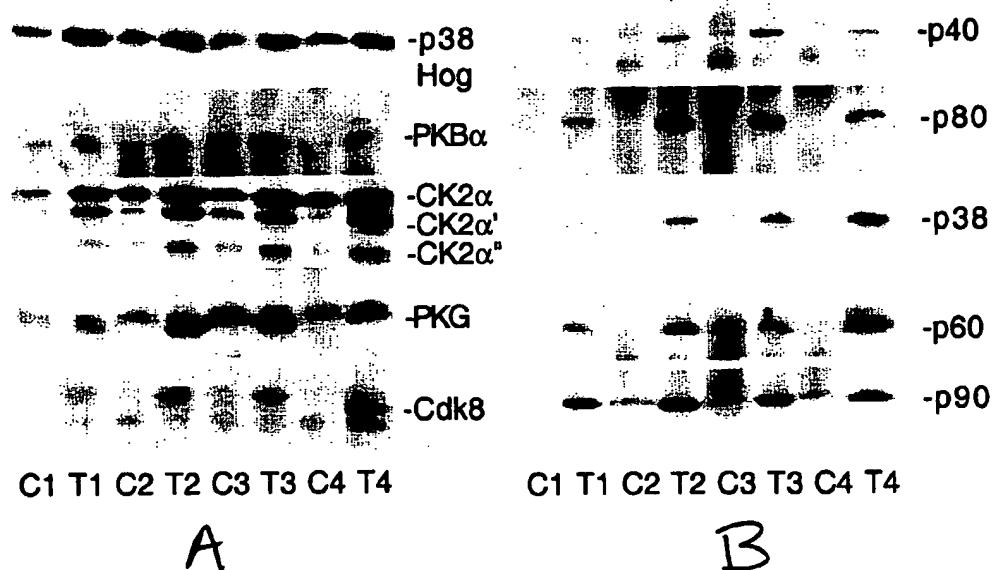


FIG. 4

FIG. 5

